

CRF Errors Corrected by the STIC System Branch

5

0590

Serial Number: 10/006,972A

CRF Processing Date: 5/28/02 05/3
 Edited by: DC
 Verified by: DC (STIC staff)

ENTERED

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: _____
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other _____
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: _____
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: _____
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: _____
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: _____
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: _____
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as _____
- ☐ Inserted mandatory headings, specifically: _____
- ☐ Corrected an obvious error in the response, specifically: _____
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: _____
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: _____
- ☐ Other: _____

*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

3/1/95



OIPE

RAW SEQUENCE LISTING

DATE: 05/28/2002

PATENT APPLICATION: US/10/006,972A

TIME: 12:35:30

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05282002\J006972A.raw

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3 <110> APPLICANT: Kenneth W. Dobie
5 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHOLIPID SCRAMBLASE 3 EXPRESSION
7 <130> FILE REFERENCE: RTS-0335
C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/006,972A
C--> 9 <141> CURRENT FILING DATE: 2001-12-04
9 <160> NUMBER OF SEQ ID NOS: 94
12 <210> SEQ ID NO: 1
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14 <212> TYPE: DNA
15 <213> ORGANISM: Artificial Sequence
17 <220> FEATURE:
18 <223> OTHER INFORMATION: Antisense Oligonucleotide
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25 <211> LENGTH: 20
26 <212> TYPE: DNA
27 <213> ORGANISM: Artificial Sequence
29 <220> FEATURE:
30 <223> OTHER INFORMATION: Antisense Oligonucleotide
32 <400> SEQUENCE: 2
33 atgcattctg cccccaagga                20
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37 <211> LENGTH: 1680
38 <212> TYPE: DNA
39 <213> ORGANISM: Homo sapiens
41 <220> FEATURE:
42 <221> NAME/KEY: CDS
43 <222> LOCATION: (144)...(1031)
45 <400> SEQUENCE: 3
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48 gtgctaggca cccgggctct tctgggggct ccagaactaa gccacccaga caccatcatc    120
50 tcgaaaaccc cagcccttct ccc atg gca ggc tac ttg ccc ccc aaa ggc tac    173
51                               Met Ala Gly Tyr Leu Pro Pro Lys Gly Tyr
52                               1           5           10
54 gcc cct tcg ccc cca cct ccc tac cct gtc acc cct ggg tac ccg gag    221
55 Ala Pro Ser Pro Pro Pro Pro Tyr Pro Val Thr Pro Gly Tyr Pro Glu
56           15           20           25
58 ccg gcg cta cat cct ggg ccc ggg cag gcg cca gtg ccc gcc cag gta    269
59 Pro Ala Leu His Pro Gly Pro Gly Gln Ala Pro Val Pro Ala Gln Val
60           30           35           40
62 cct gcc cca gct ccc ggc ttc gcc ctc ttc ccc tcg cct ggc ccc gtg    317
63 Pro Ala Pro Ala Pro Gly Phe Ala Leu Phe Pro Ser Pro Gly Pro Val

```

RAW SEQUENCE LISTING

DATE: 05/28/2002

PATENT APPLICATION: US/10/006,972A

TIME: 12:35:30

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05282002\J006972A.raw

64	45	50	55	
66	gcc ttg ggg tct gct gcc ccc ttc ttg cca ctg cca ggg gtg cct tct	365		
67	Ala Leu Gly Ser Ala Ala Pro Phe Leu Pro Leu Pro Gly Val Pro Ser			
68	60	65	70	
70	ggc ctc gaa ttc ctg gtg cag att gat cag att ttg att cac cag aag	413		
71	Gly Leu Glu Phe Leu Val Gln Ile Asp Gln Ile Leu Ile His Gln Lys			
72	75	80	85	90
74	gct gag cga gtg gaa acg ttc cta ggc tgg gag acc tgt aat cgg tat	461		
75	Ala Glu Arg Val Glu Thr Phe Leu Gly Trp Glu Thr Cys Asn Arg Tyr			
76	95	100	105	
78	gaa ctg cgc tct ggg gcc ggg cag ccc ctg ggt cag gcg gcc gag gag	509		
79	Glu Leu Arg Ser Gly Ala Gly Gln Pro Leu Gly Gln Ala Ala Glu Glu			
80	110	115	120	
82	agc aac tgc tgc gcc cgt ctg tgc tgt ggc gcc cgc cgg ccg ctg cgt	557		
83	Ser Asn Cys Cys Ala Arg Leu Cys Cys Gly Ala Arg Arg Pro Leu Arg			
84	125	130	135	
86	gtc cgc ctg gcc gac ccc ggg gac cgt gag gtg ctg cgt ttg ctc cgc	605		
87	Val Arg Leu Ala Asp Pro Gly Asp Arg Glu Val Leu Arg Leu Leu Arg			
88	140	145	150	
90	ccg ctg cac tgt ggc tgc agc tgc tgc ccc tgt ggc ctc cag gag atg	653		
91	Pro Leu His Cys Gly Cys Ser Cys Cys Pro Cys Gly Leu Gln Glu Met			
92	155	160	165	170
94	gaa gta cag gct cca cca ggc acc acc att ggc cac gtg cta cag acc	701		
95	Glu Val Gln Ala Pro Pro Gly Thr Thr Ile Gly His Val Leu Gln Thr			
96	175	180	185	
98	tgg cat ccc ttc ctc ccc aag ttc tcc atc cag gat gcc gat cgc cag	749		
99	Trp His Pro Phe Leu Pro Lys Phe Ser Ile Gln Asp Ala Asp Arg Gln			
100	190	195	200	
102	aca gtc ttg cga gtg gtg ggg ccc tgc tgg acc tgt ggc tgt ggc aca	797		
103	Thr Val Leu Arg Val Val Gly Pro Cys Trp Thr Cys Gly Cys Gly Thr			
104	205	210	215	
106	gac acc aac ttt gag gtg aag act cgg gat gaa tcc cgc agt gtg ggc	845		
107	Asp Thr Asn Phe Glu Val Lys Thr Arg Asp Glu Ser Arg Ser Val Gly			
108	220	225	230	
110	cgc atc agc aag cag tgg ggg ggc ctg gtc cga gaa gcc ctc aca gat	893		
111	Arg Ile Ser Lys Gln Trp Gly Gly Leu Val Arg Glu Ala Leu Thr Asp			
112	235	240	245	250
114	gca gat gac ttt ggc cta cag ttc ccg ctg gac ctg gat gtg agg gtg	941		
115	Ala Asp Asp Phe Gly Leu Gln Phe Pro Leu Asp Leu Asp Val Arg Val			
116	255	260	265	
118	aag gct gtg ctg ctg gga gcc aca ttc ctc att gac tac atg ttc ttt	989		
119	Lys Ala Val Leu Leu Gly Ala Thr Phe Leu Ile Asp Tyr Met Phe Phe			
120	270	275	280	
122	gag aag cga gga ggc gct ggg ccc tct gcc atc acc agt tag aggccaccat	1041		
123	Glu Lys Arg Gly Gly Ala Gly Pro Ser Ala Ile Thr Ser			
124	285	290	295	
126	ggtgtgagga gaccatcacc tcgaccagaa ctccagatgg tcacctgccc tggcccctcc	1101		
128	tctgggcagc ccttttctc catgtacact gcaggggaca gaaggggggc cccatcccta	1161		
130	ccctactccc tggccgcctg cccctgtggt tcccaaggag gggtatgtat gagagccgct	1221		

RAW SEQUENCE LISTING

DATE: 05/28/2002

PATENT APPLICATION: US/10/006,972A

TIME: 12:35:30

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05282002\J006972A.raw

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132 ctccctgctac ctcccaccac tgtcccagca gtccctcggc acacaggcat atcagctttc 1281
134 acacttttccc catgcaactct ctcccacccc ctccagggc ctctgctcca aaggaggcct 1341
136 ctggaaccca ggactctggg gttttacaag agggctggg tgtggaagg caagctgcac 1401
138 caaagacggt ggatatagcc accgcccccc cgcgctgcc tagcatctgc ttggccaatt 1461
140 agttcagcct cagaccatgg cactttgagg gggctctctac ctccccatca acagctgcag 1521
142 ggggacccca gtgccaactt cctctccac tagggccctg ccttcagctg gtgcttgctg 1581
144 cgattcctgt gccttatgta actgcccttc cttcccttgc cctaggaaaa aggctgcac 1641
146 tttatatgtt acattcatat aaactttgta actttttgg 1680
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150 <211> LENGTH: 20
151 <212> TYPE: DNA
152 <213> ORGANISM: Artificial Sequence
154 <220> FEATURE:
155 <223> OTHER INFORMATION: PCR Primer
157 <400> SEQUENCE: 4
158 gtccgagaag ccctcacaga 20
161 <210> SEQ ID NO: 5
162 <211> LENGTH: 19
163 <212> TYPE: DNA
164 <213> ORGANISM: Artificial Sequence
166 <220> FEATURE:
167 <223> OTHER INFORMATION: PCR Primer
169 <400> SEQUENCE: 5
170 gccttcaccc tcacatcca 19
173 <210> SEQ ID NO: 6
174 <211> LENGTH: 27
175 <212> TYPE: DNA
176 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: PCR Probe
181 <400> SEQUENCE: 6
182 cagatgactt tggcctacag ttccccg 27
185 <210> SEQ ID NO: 7
186 <211> LENGTH: 19
187 <212> TYPE: DNA
188 <213> ORGANISM: Artificial Sequence
190 <220> FEATURE:
191 <223> OTHER INFORMATION: PCR Primer
193 <400> SEQUENCE: 7
194 gaaggtgaag gtcggagtc 19
197 <210> SEQ ID NO: 8
198 <211> LENGTH: 20
199 <212> TYPE: DNA
200 <213> ORGANISM: Artificial Sequence
202 <220> FEATURE:
203 <223> OTHER INFORMATION: PCR Primer
205 <400> SEQUENCE: 8
206 gaagatggtg atgggatttc 20
209 <210> SEQ ID NO: 9

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RAW SEQUENCE LISTING

DATE: 05/28/2002

PATENT APPLICATION: US/10/006,972A

TIME: 12:35:30

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05282002\J006972A.raw

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210 <211> LENGTH: 20
211 <212> TYPE: DNA
212 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: PCR Probe
217 <400> SEQUENCE: 9
218 caagcttccc gttctcagcc 20
221 <210> SEQ ID NO: 10
222 <211> LENGTH: 596
223 <212> TYPE: DNA
224 <213> ORGANISM: Homo sapiens
226 <220> FEATURE:
W--> 227 <221> NAME/KEY: exon:exon junction
228 <222> LOCATION: (333)...(334)
229 <223> OTHER INFORMATION: exon 5:exon 6b
W--> 231 <221> NAME/KEY: exon:exon junction
232 <222> LOCATION: (423)...(424)
233 <223> OTHER INFORMATION: exon 6b:exon 7
W--> 236 <400> 10
237 ttgggggtctg ctgccccctt cttgccactg ccagggtgcc ttctggcctc gaattcctgg 60
239 tgcagattga tcagattttg attcaccaga aggctgagcg agtggaacg ttcctagtgc 120
241 tgggagacct gtaatcggta tgaactgcgc tctggggcct gggcagcccc tgggtcaggc 180
243 ggccgaggag agcaactgct gcgcccgtct gtgctgtggc tgcccgccgg cctgctgcgt 240
245 gtccgcctgg ccgaccccg ggaccgtgag gtgctgcgtt tgctccgccc gctgcactgt 300
247 ggctgcagct gctgcccctg tggcctccag gagttctcca tccaggatgc cgatcgccag 360
249 acagtcttgc gagtgggtgg gccctgctgg acctgtggct gtggcacaga caccaacttt 420
251 gaggtgaaga ctcgggatga atcccgagc gtgggcccga tcagcaagca gtgtgggggg 480
253 cctggtccga gaagccctca cagatgcaga tgactttggc ctacagttcc cgctggacct 540
255 ggatgtgagg gtgaaggctg tgctgctggg agccacattc ctcatttgac tactgt 596
258 <210> SEQ ID NO: 11
W--> 259 <400> SEQUENCE: 11
W--> 260 000
262 <210> SEQ ID NO: 12
W--> 263 <400> SEQUENCE: 12
W--> 264 000
266 <210> SEQ ID NO: 13
W--> 267 <400> SEQUENCE: 13
W--> 268 000
270 <210> SEQ ID NO: 14
271 <211> LENGTH: 20
272 <212> TYPE: DNA
273 <213> ORGANISM: Artificial Sequence
275 <220> FEATURE:
276 <223> OTHER INFORMATION: Antisense Oligonucleotide
278 <400> SEQUENCE: 14
279 cggcagctca caggagctgg 20
282 <210> SEQ ID NO: 15
283 <211> LENGTH: 20
284 <212> TYPE: DNA

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RAW SEQUENCE LISTING

DATE: 05/28/2002

PATENT APPLICATION: US/10/006,972A

TIME: 12:35:30

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05282002\J006972A.raw

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285 <213> ORGANISM: Artificial Sequence
287 <220> FEATURE:
288 <223> OTHER INFORMATION: Antisense Oligonucleotide
290 <400> SEQUENCE: 15
291 gcactcggca gctcacagga                                20
294 <210> SEQ ID NO: 16
295 <211> LENGTH: 20
296 <212> TYPE: DNA
297 <213> ORGANISM: Artificial Sequence
299 <220> FEATURE:
300 <223> OTHER INFORMATION: Antisense Oligonucleotide
302 <400> SEQUENCE: 16
303 tgcctagcac tcggcagctc                                20
306 <210> SEQ ID NO: 17
307 <211> LENGTH: 20
308 <212> TYPE: DNA
309 <213> ORGANISM: Artificial Sequence
311 <220> FEATURE:
312 <223> OTHER INFORMATION: Antisense Oligonucleotide
314 <400> SEQUENCE: 17
315 tggcttagtt ctggagcccc                                20
318 <210> SEQ ID NO: 18
319 <211> LENGTH: 20
320 <212> TYPE: DNA
321 <213> ORGANISM: Artificial Sequence
323 <220> FEATURE:
324 <223> OTHER INFORMATION: Antisense Oligonucleotide
326 <400> SEQUENCE: 18
327 tgggtgtctgg gtggcttagt                                20
330 <210> SEQ ID NO: 19
331 <211> LENGTH: 20
332 <212> TYPE: DNA
333 <213> ORGANISM: Artificial Sequence
335 <220> FEATURE:
336 <223> OTHER INFORMATION: Antisense Oligonucleotide
338 <400> SEQUENCE: 19
339 tcgagatgat ggtgtctggg                                20
342 <210> SEQ ID NO: 20
343 <211> LENGTH: 20
344 <212> TYPE: DNA
345 <213> ORGANISM: Artificial Sequence
347 <220> FEATURE:
348 <223> OTHER INFORMATION: Antisense Oligonucleotide
350 <400> SEQUENCE: 20
351 gcaagtagcc tgccatggga                                20
354 <210> SEQ ID NO: 21
355 <211> LENGTH: 20
356 <212> TYPE: DNA
357 <213> ORGANISM: Artificial Sequence

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VERIFICATION SUMMARY

DATE: 05/28/2002

PATENT APPLICATION: US/10/006,972A

TIME: 12:35:31

Input Set : A:\PTO.DC.txt

Output Set: N:\CRF3\05282002\J006972A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:227 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:231 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:236 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:10
L:259 M:283 W: Missing Blank Line separator, <400> field identifier
L:260 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (11) SEQUENCE:
L:263 M:283 W: Missing Blank Line separator, <400> field identifier
L:264 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (12) SEQUENCE:
L:267 M:283 W: Missing Blank Line separator, <400> field identifier
L:268 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (13) SEQUENCE:
L:1087 M:283 W: Missing Blank Line separator, <400> field identifier
L:1088 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (82) SEQUENCE:
L:1092 M:283 W: Missing Blank Line separator, <400> field identifier
L:1093 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (83) SEQUENCE:
L:1097 M:283 W: Missing Blank Line separator, <400> field identifier
L:1098 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (84) SEQUENCE:
L:1102 M:283 W: Missing Blank Line separator, <400> field identifier
L:1103 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (85) SEQUENCE:
L:1107 M:283 W: Missing Blank Line separator, <400> field identifier
L:1108 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (86) SEQUENCE:
L:1136 M:283 W: Missing Blank Line separator, <400> field identifier
L:1137 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (89) SEQUENCE:
L:1141 M:283 W: Missing Blank Line separator, <400> field identifier
L:1142 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (90) SEQUENCE:
L:1146 M:283 W: Missing Blank Line separator, <400> field identifier
L:1147 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (91) SEQUENCE:



Does Not Comply
Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/006,972A

DATE: 05/17/2002

TIME: 14:20:55

Input Set : A:\RTS-335_Seq_ASCII.txt

Output Set: N:\CRF3\05172002\J006972A.raw

3 <110> APPLICANT: Kenneth W. Dobie
 5 <120> TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHOLIPID SCRAMBLASE 3 EXPRESSION
 7 <130> FILE REFERENCE: RTS-0335
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/006,972A
 C--> 9 <141> CURRENT FILING DATE: 2001-12-04
 9 <160> NUMBER OF SEQ ID NOS: 94

ERRORED SEQUENCES

1186 <210> SEQ ID NO: 94
 1187 <211> LENGTH: 546
 1188 <212> TYPE: DNA
 1189 <213> ORGANISM: Homo sapiens
 1191 <400> SEQUENCE: 94
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 1193 gctccagtca gaggcgccgc ccaagagacc ctgggccggc gccgggcgca gctgcctctc 120
 1194 cgtctttgtg tctgtctctg tgtctgtctg gctatctccg agtttgctc cgttccaga 180
 1195 actaagccac ccagacacca tcctctcgaa aaccccgacc ctctcccat ggcaggctac 240
 1196 ttgcccccca aaggctacgc cccttcgccc ccacctccct accctgtcac ccctgggtac 300
 1197 ccgtgcgtg tccgcctggc cgaccccggg gaccgtgagg tgctgcgttt gctccgccc 360
 1198 gtgcactggg ggcttcgagg tgtgttgccc tttggggcct ccaggagatg gatgtacggg 420
 1199 ctccaccagg caccacctat gggccacgtg ctacagacct ggcacccctt cctcccaaag 480
 1200 ttctccatcc aggatgccga tcgccagaca gtcttgcgaa gtgggtggggc cctgcctgga 540
 1201 cctgtg 546
 E--> 1206 ① - delete

VERIFICATION SUMMARY

DATE: 05/17/2002

PATENT APPLICATION: US/10/006,972A

TIME: 14:20:56

Input Set : A:\RTS-335_Seq_ASCII.txt

Output Set: N:\CRF3\05172002\J006972A.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:227 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:231 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:10
L:236 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:10
L:259 M:283 W: Missing Blank Line separator, <400> field identifier
L:260 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (11) SEQUENCE:
L:263 M:283 W: Missing Blank Line separator, <400> field identifier
L:264 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (12) SEQUENCE:
L:267 M:283 W: Missing Blank Line separator, <400> field identifier
L:268 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (13) SEQUENCE:
L:1087 M:283 W: Missing Blank Line separator, <400> field identifier
L:1088 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (82) SEQUENCE:
L:1092 M:283 W: Missing Blank Line separator, <400> field identifier
L:1093 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (83) SEQUENCE:
L:1097 M:283 W: Missing Blank Line separator, <400> field identifier
L:1098 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (84) SEQUENCE:
L:1102 M:283 W: Missing Blank Line separator, <400> field identifier
L:1103 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (85) SEQUENCE:
L:1107 M:283 W: Missing Blank Line separator, <400> field identifier
L:1108 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (86) SEQUENCE:
L:1136 M:283 W: Missing Blank Line separator, <400> field identifier
L:1137 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (89) SEQUENCE:
L:1141 M:283 W: Missing Blank Line separator, <400> field identifier
L:1142 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (90) SEQUENCE:
L:1146 M:283 W: Missing Blank Line separator, <400> field identifier
L:1147 M:300 W: (50) Intentionally skipped Sequence, : Sequence Id (91) SEQUENCE:
L:1206 M:254 E: No. of Bases conflict, this line has no nucleotides.